

We claim:

1. A method for determining a profile data set for a subject, based on a sample from the subject, the sample providing a source of RNAs, the method comprising:  
quantitatively measuring an amount of RNA corresponding to each of at least  
5 two constituents from any one of Tables 1 through 8, where the profile data set comprises the measure of each constituent and wherein such measure for each constituent is obtained under measurement conditions that are substantially reproducible.
2. A method according to claim 1, wherein the measurement conditions that are  
10 reproducible include intra-assay variability and inter-assay variability.
3. A method according to claim 2, wherein intra-assay variability is reproducible such that the average coefficient of variation for repeated measurements for each constituent from the same sample is less than 1 percent.
4. A method according to claim 2, wherein inter-assay variability is reproducible  
15 such that the average coefficient of variation for measurements for each constituent in distinct samples of the same material is less than 2 percent.
5. A method according to claim 1, wherein the profile data set is informative about a biological condition for the subject with respect to the circumstances of the subject at the time the sample was obtained.
- 20 6. A method according to claim 5, wherein the profile data set provides guidance for medical or surgical intervention by allopathic or naturpathic means.

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